SEQUENCE LISTING

<110> MAIER, THOMAS

<120> METHOD FOR FERMENTATIVE PRODUCTION OF AMINO ACIDS AND AMINO ACID DERIVATIVES OF THE PHOSPHOGLYCERATE FAMILY

<130> MAIER, T-2

<150> GERMAN NO. 102 32 930.3

<151> 2002-07-19

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<170> PatentIn version 3.1

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<212> DNA

<213> Escherichia coli

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                                                      Val Thr Pro
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acc ctt tta agt gct ttt tgg act tac acc ctg att acc gct atg acg      166
Thr Leu Leu Ser Ala Phe Trp Thr Tyr Thr Leu Ile Thr Ala Met Thr
    5                   10                  15

cca gga ccg aac aat att ctc gcc ctt agc tct gct acg tcg cat gga      214
Pro Gly Pro Asn Asn Ile Leu Ala Leu Ser Ser Ala Thr Ser His Gly
20                  25                  30                  35

ttt cgt caa agt acc cgc gtg ctg gca ggg atg agt ctg gga ttt ttg      262
Phe Arg Gln Ser Thr Arg Val Leu Ala Gly Met Ser Leu Gly Phe Leu
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att gtg atg tta ctg tgt gcg ggc att tca ttt tca ctg gca gtg att      310
Ile Val Met Leu Leu Cys Ala Gly Ile Ser Phe Ser Leu Ala Val Ile
            55                  60                  65

gac ccg gca gcg gta cac ctt ttg agt tgg gcg ggg gcg gca tat att      358
Asp Pro Ala Ala Val His Leu Leu Ser Trp Ala Gly Ala Ala Tyr Ile
        70                  75                  80

gtc tgg ctg gcg tgg aaa atc gcc acc agc cca aca aag gaa gac gga      406
Val Trp Leu Ala Trp Lys Ile Ala Thr Ser Pro Thr Lys Glu Asp Gly
    85                  90                  95

ctt cag gca aaa cca atc agc ttt tgg gcc agc ttt gct ttg cag ttt      454
Leu Gln Ala Lys Pro Ile Ser Phe Trp Ala Ser Phe Ala Leu Gln Phe
100                 105                 110                 115

gtg aac gtc aaa atc att ttg tac ggt gtt acg gca ctg tcg acg ttt      502
Val Asn Val Lys Ile Ile Leu Tyr Gly Val Thr Ala Leu Ser Thr Phe
                120                 125                 130

gtt ctg ccg caa aca cag gcg tta agc tgg gta gtt ggc gtc agc gtt      550
Val Leu Pro Gln Thr Gln Ala Leu Ser Trp Val Val Gly Val Ser Val
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ggg cat ctg ttt cag cga ttg ttt cgc cag tat ggt cgc cag tta aat 646
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165 170 175

atc gtg ctt gcc ctg ttg ctg gtc tat tgc gcg gta cgc att ttc tat 694
Ile Val Leu Ala Leu Leu Leu Val Tyr Cys Ala Val Arg Ile Phe Tyr
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Ser His Gly Phe Arg Gln Ser Thr Arg Val Leu Ala Gly Met Ser Leu
35 40 45

Gly Phe Leu Ile Val Met Leu Leu Cys Ala Gly Ile Ser Phe Ser Leu
50 55 60

Ala Val Ile Asp Pro Ala Ala Val His Leu Leu Ser Trp Ala Gly Ala
65 70 75 80

Ala Tyr Ile Val Trp Leu Ala Trp Lys Ile Ala Thr Ser Pro Thr Lys
85 90 95

Glu Asp Gly Leu Gln Ala Lys Pro Ile Ser Phe Trp Ala Ser Phe Ala
100 105 110

Leu Gln Phe Val Asn Val Lys Ile Ile Leu Tyr Gly Val Thr Ala Leu
115 120 125

Ser Thr Phe Val Leu Pro Gln Thr Gln Ala Leu Ser Trp Val Val Gly
130 135 140

Val Ser Val Leu Leu Ala Met Ile Gly Thr Phe Gly Asn Val Cys Trp
145 150 155 160

Ala Leu Ala Gly His Leu Phe Gln Arg Leu Phe Arg Gln Tyr Gly Arg
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Gln Leu Asn Ile Val Leu Ala Leu Leu Leu Val Tyr Cys Ala Val Arg
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Ile Phe Tyr
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